



SEQUENCE LISTING

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Thompson, Leslie M.
Housman, David E.

<120> INHIBITION OF PROTEIN-PROTEIN INTERACTION

<130> 01997-289001

<140> US 09/933,638

<141> 2001-08-20

<150> US 60/226,502

<151> 2000-08-18

<160> 12

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Val Gln Ile Val Tyr Lys

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 Phe Arg Lys

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 Arg Phe Ala Ala Val Ile Met Arg Ile Arg Glu Pro Arg Thr Thr Ala
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 Leu Ile Phe Ser Ser Gly Lys Met Val Cys Thr Gly Ala Lys Ser Glu
 35 40 45
 Glu Gln Ser Arg Leu Ala Ala Arg Lys Tyr Ala Arg
 50 55 60

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 Glu Glu Gln Ser Arg Leu Ala Ala Arg Lys Tyr Ala Arg Val Val Gln
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 Lys Leu Gly Phe Pro Ala Lys Phe Leu Asp Phe Lys Ile Gln Asn Met
 20 25 30

Val Gly Ser Cys Asp Val Lys Phe Pro Ile Arg Leu Glu Gly Leu Val
 35 40 45
 Leu Thr His Gln Gln Phe
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<220>
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 Arg Leu Glu Gly Leu Val Leu Thr His Gln Gln Phe Ser Ser Tyr Glu
 1 5 10 15
 Pro Glu Leu Phe Pro Gly Leu Ile Tyr Arg Met Ile Lys Pro Arg Ile
 20 25 30
 Val Leu Leu Ile Phe Val Ser Gly Lys Val Val Leu Thr Gly Ala Lys
 35 40 45
 Val Arg Ala Glu Ile Tyr Glu Ala Phe Glu Asn Ile Tyr Pro Ile Leu
 50 55 60
 Lys Gly Phe Arg Lys
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<220>
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<400> 11

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Leu Glu Gly Leu Val Leu Thr His Gln Gln Phe Ser Ser Tyr Glu Pro
 1           5           10           15
Glu Leu Phe Pro Gly Leu Ile Tyr Arg Met Ile Lys Pro Arg Ile Val
           20           25           30
Leu Leu Ile Phe Val Ser Gly Lys Val Val Leu Thr Gly Ala Lys Val
           35           40           45
Arg Ala Glu Ile Tyr Glu Ala Phe Glu Asn Ile Tyr Pro Ile Leu Lys
           50           55           60
Gly Phe Arg Lys
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<213> Homo sapiens

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           20           25           30
Pro Tyr Gly Thr Gly Leu Thr Pro Gln Pro Ile Gln Asn Thr Asn Ser
           35           40           45
Leu Ser Ile Leu Glu Glu Gln Gln Arg Gln Gln Gln Gln Gln Gln
           50           55           60
Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln
           65           70           75           80
Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Gln Ala
           85           90           95
Val Ala Ala Ala Ala Val Gln Gln Ser Thr Ser Gln Gln Ala Thr Gln
           100          105          110
Gly Thr Ser Gly Gln Ala Pro Gln Leu Phe His Ser Gln Thr Leu Thr
           115          120          125
Thr Ala Pro Leu Pro Gly Thr Thr Pro Leu Tyr Pro Ser Pro Met Thr
           130          135          140
Pro Met Thr Pro Ile Thr Pro Ala Thr Pro Ala Ser Glu Ser Ser Gly
           145          150          155          160
Ile Val Pro Gln Leu Gln Asn Ile Val Ser Thr Val Asn Leu Gly Cys
           165          170          175
Lys Leu Asp Leu Lys Thr Ile Ala Leu Arg Ala Arg Asn Ala Glu Tyr
           180          185          190
Asn Pro Lys Arg Phe Ala Ala Val Ile Met Arg Ile Arg Glu Pro Arg
           195          200          205
Thr Thr Ala Leu Ile Phe Ser Ser Gly Lys Met Val Cys Thr Gly Ala
           210          215          220
Lys Glu Glu Gln Ser Arg Leu Ala Ala Arg Lys Tyr Ala Arg Val Val
           225          230          235          240
Gln Lys Leu Gly Phe Pro Ala Lys Phe Leu Asp Phe Lys Ile Gln Asn
           245          250          255
Met Val Gly Ser Cys Asp Val Lys Phe Pro Ile Arg Leu Glu Gly Leu
           260          265          270
Val Leu Thr His Gln Gln Phe Ser Ser Tyr Glu Pro Glu Leu Phe Pro
           275          280          285
Gly Leu Ile Tyr Arg Met Ile Lys Pro Arg Ile Val Leu Leu Ile Phe
           290          295          300

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[illegible]